

Fig. 1

distal UP

nnAAA (A/T) (A/T) T (A/T) TTTTnnAAAAnnn
proximal UP

Fig. 2A

-66 -59 UP element -38
CCCCGTCAGAAAATTATTTTAAATTCCTCTTGTCAGGCCCGAATAACTCCCTAT
AATGCGCCAC

+1 +50
CACTGACACGGAACAACGGCAAACACGCCGCCGGGTCAGCGGGGTTCCTCCT

Fig. 2B

•Direct targeting

MEF	C (TTAAAAATAA) C	SEQ: 22
780BP	(TTGAAAAATCAA) CGCT	SEQ: 23

•Overlapping Targeting (test for up or down-stream)

UL9	(ttttTGTT) CGCAC (TTtttttttt)	SEQ: 24
NFκB	(tttttGGG[AtTTT] CQttttt)	SEQ: 25
LacO	(aaaaAATT) GTGAGCGCTCAC (AATTtttt)	SEQ: 26
NtBBF1	(tttACT[TTA)tttt]	

Fig. 3

rrnB P1 promoter UP Sequences

RLG3097 (core)	GACTGCAGTGGTACCTAGGAGG	SEQ: 14
RLG3074 (wt)	AGAAAATTATTTTAAATTCCT	SEQ: 13
RLG4192	GGAAAATTTTTTTTCAAAAGTA	SEQ: 16
RLG4174	TGAAATTTATTTTGCGAAAGGG	SEQ: 17

Fig. 4A

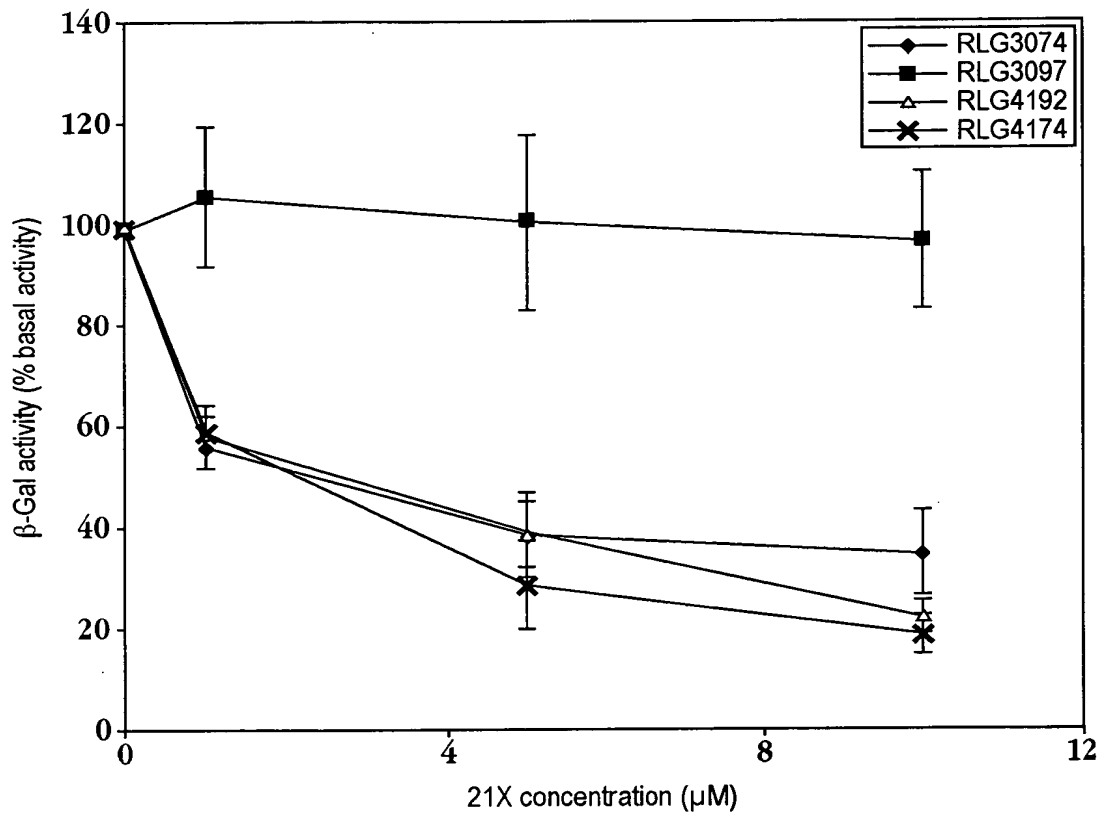


Fig. 4B

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YK 202LX (52-mer) 5' CATGGACG CCACTG AGCCG TTTT TGTTCCGCACTT GAGGCGAGTCGATGCACC 3'
                  3' GTACCTGC GGTGAC TCGCAAAA ACAAGCGTGAA CTCGCTCAGCTACGTGG 5'

YK 202RX-A (54-mer) 5' CATGGACG CCACTG AGCCG TGTTCCGCACTT TTTTGTGAGGCGAGTCGATGCACC 3'
YK 202RX-B (54-mer) 3' GTACCTGC GGTGAC TCGGC ACAAGCGTGAA AAAAAACTCCGCTCAGCTACGTGGB 5'

YK 202LRX (58-mer) 5' CATGGACG CCACTG AGCCG TTTT TGTTCCGCACTT TTTTGTGAGGCGAGTCGATGCACC 3'
                  3' GTACCTGC GGTGAC TCGCAAAA ACAAGCGTGAA AAAAAACTCCGCTCAGCTACGTGG 5'

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Fig. 5

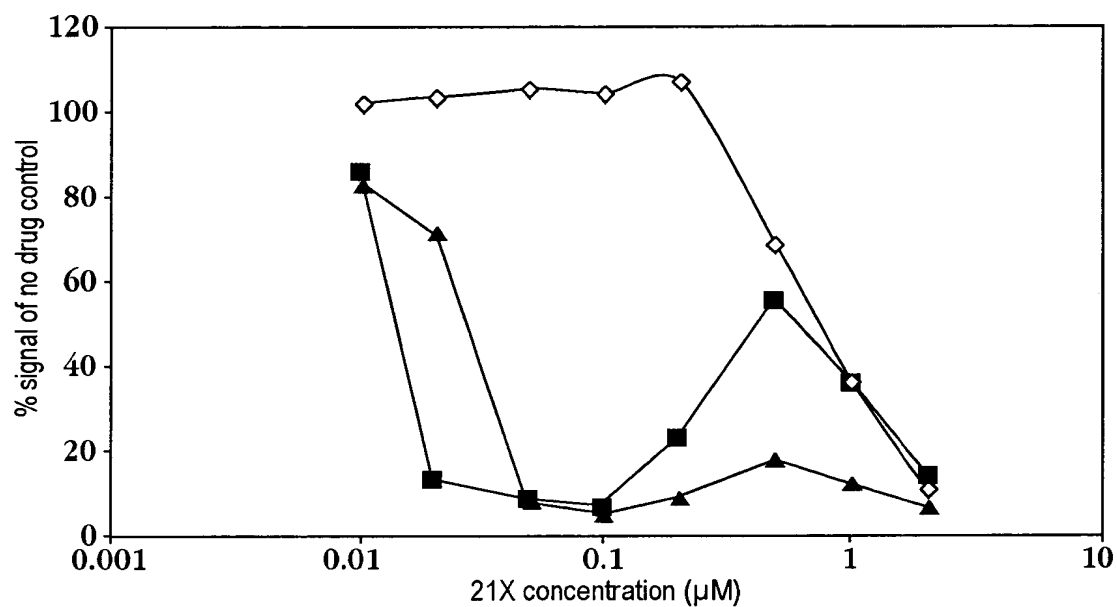


Fig. 6

JF 101 (NFKB1) (50mer) (right side)

5' cgac cgtgctcgag **TTAACGGGACTTTCCAAaaa** cgatcg gact ggactc 3'
3' gctg gcacgagctc **AATTGCCCTGAAAGGTTttt** gctagc ctga cctgag 5'

JF 102 (NFKB2) (60mer) (right side)

5' cgac cgtgctcgag **TTAACGGGAtTTTCCAAaaa** cgatcg gact ggactc 3'
3' gctg gcacgagctc **AATTGCCCTaAAAGGTTttt** gctagc ctga cctgag 5'

JF 103 (NFKB3) (60mer) (both side)

5' cgac cgtgctcgag **aaattGGGAtTTTCCAAaaa** cgatcg gact ggactc 3'
3' gctg gcacgagctc **tttaaCCCTaAAAGGTTttt** gctagc ctga cctgag 5'

Fig. 7

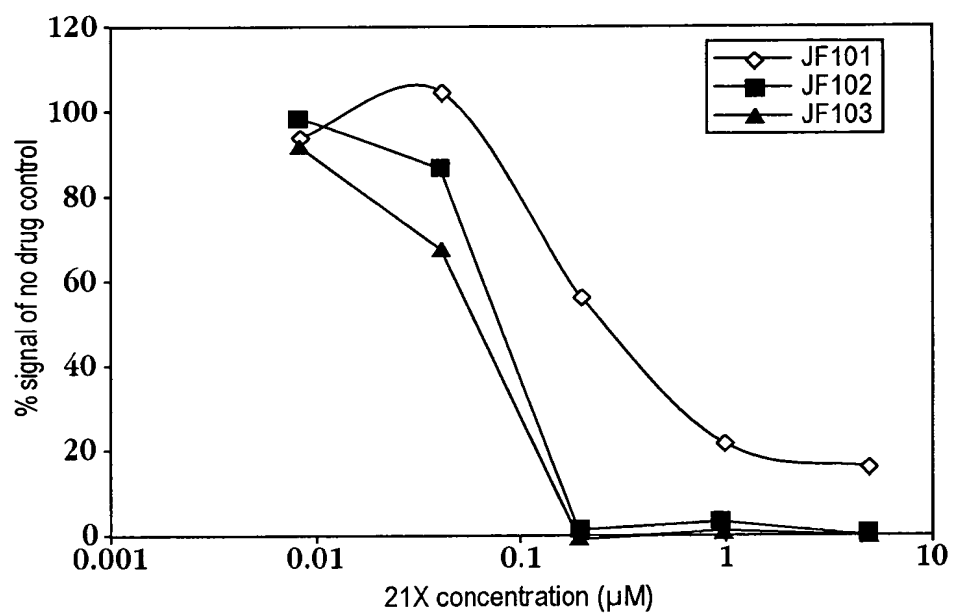


Fig. 8A

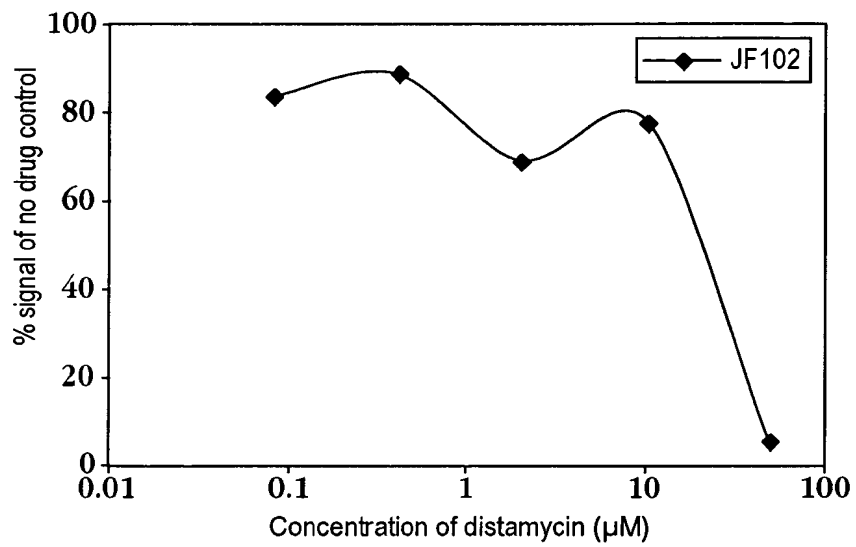


Fig. 8B

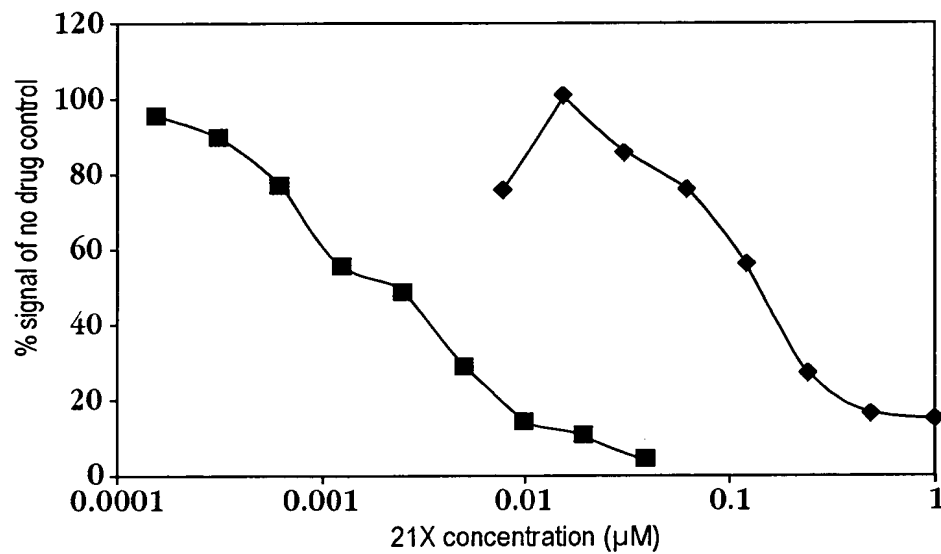


Fig. 9

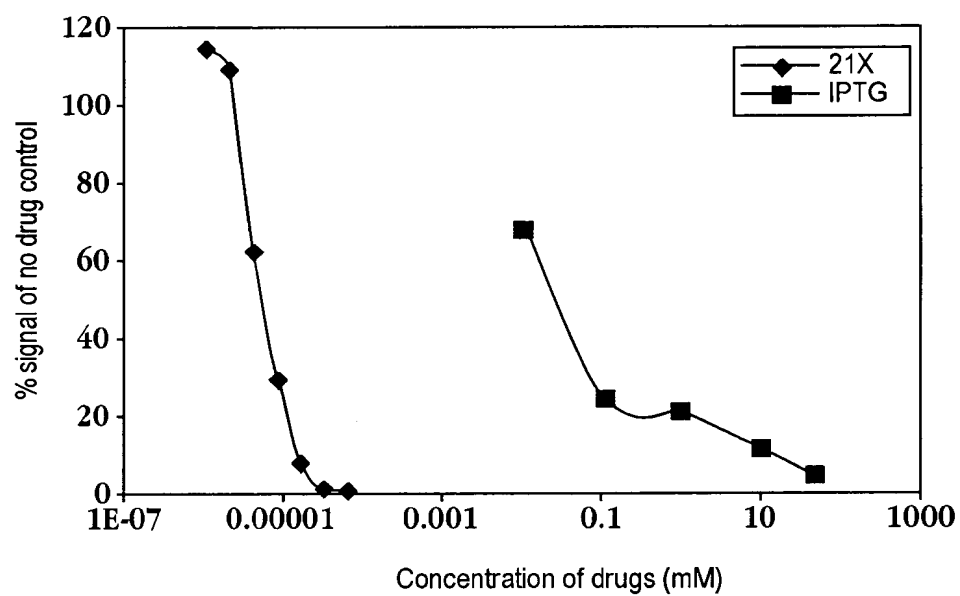


Fig. 10

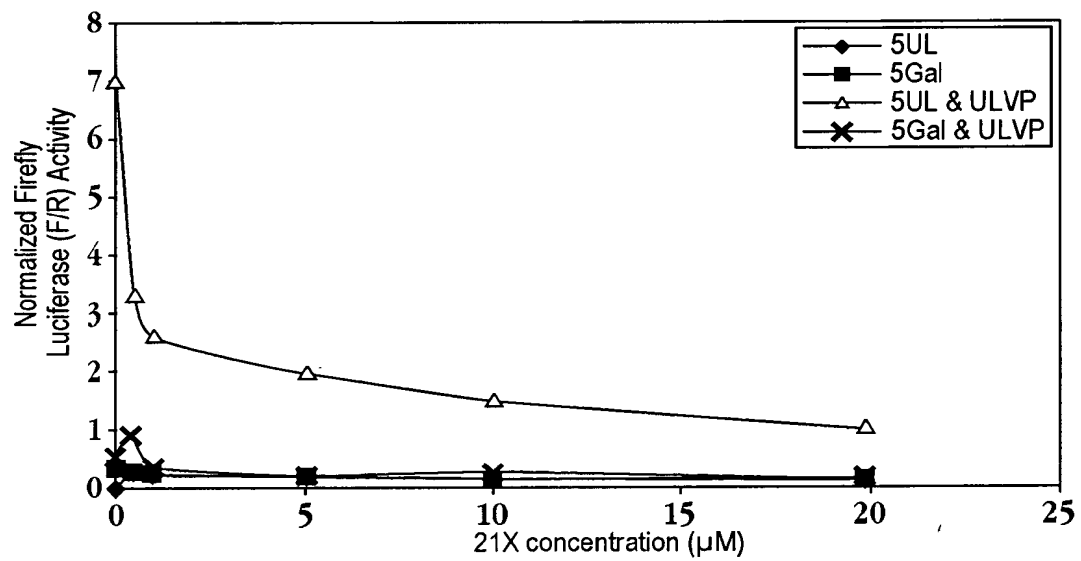


Fig. 11

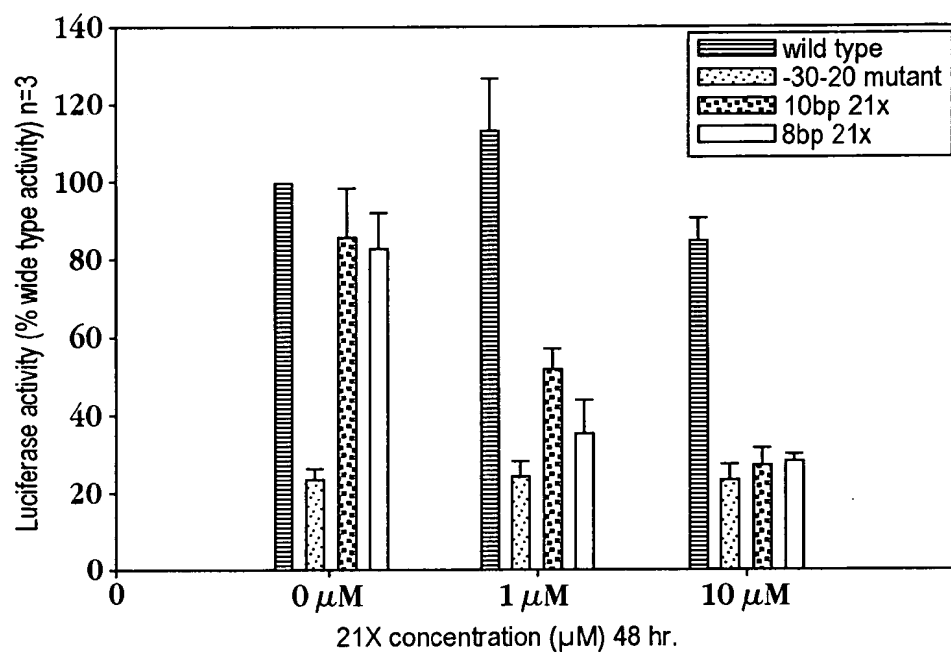


Fig. 12

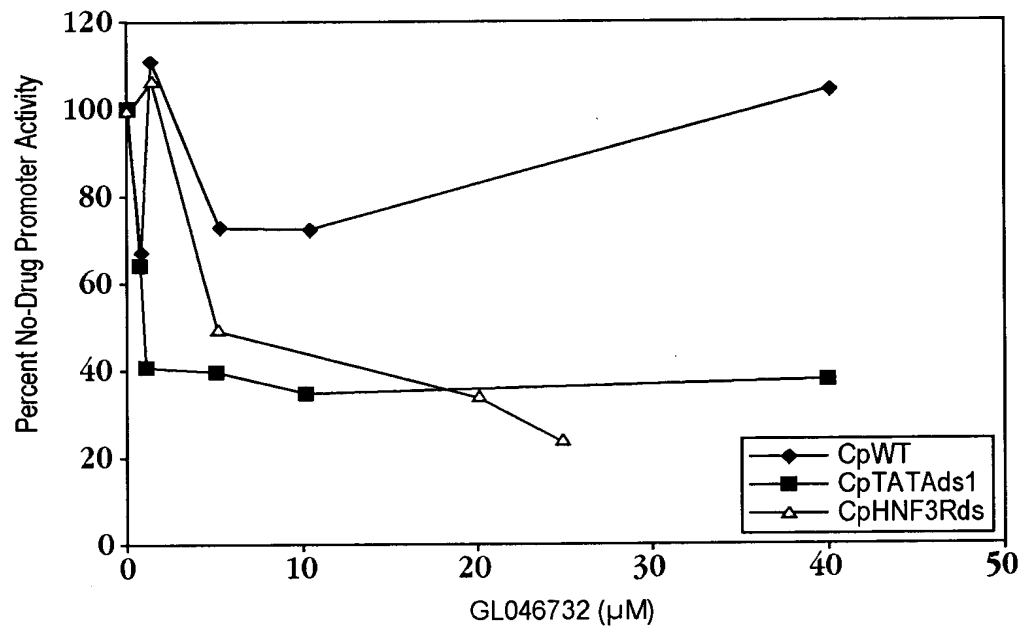


Fig. 13

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CCCATATATGGAGTTCGCGGTTACATAACTTACGGTAAATGGCCCGCCTGGCTGACCGCCCAACG
ACCCCGCCCATTTGACGTCAATAATGACGTATGTTCCCATAGTAACGCCAATAGGGACTTTCCAT
TGACGTCAATGGGTGGAGTATTTACGGTAAACTGCCCACTTGGCAGTACATCAAGTGTATCATAT
GCCAAGTCCGCCCCCTATTGACGTCAATGACGGTAAATGGCCCGCCTGGCATTATGCCAGTACA
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AGGCAGAAGTATGCAAAGCATGCATCTCAATTAGTCAGCAACCATAGTCCCGCCCCCTAACTCCGC

Fig. 14A

CCATCCCGCCCCCTAACTCCGCCCAGTTCGCCCCATTCTCCGCCCCATGGCTGACTAATTTTTTTTT
ATTTATGCAGAGGCCGAGGCCGCTCGGCCTCTGAGCTATTCCAGAAGTAGTGAGGAGGCTTTTT
TGGAGGCCTAGGCTTTTGC AAAAAGCTTGATTCTTCTGACACAACAGTCTCGAACTTAAGGCTAG
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GGAAAACGCCAGCAACGCGGCCTTTTACGGTTCTTGGCCTTTTGCTGGCCTTTTGCTCACATG
GCTCGACAGATCT

Fig. 14B

TCAATATTGGCCATTAGCCATATTATTCATTGGTTATATAGCATAAATCAATATTGGCTATTGGC
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CCATGTTGGCATTGATTATTGACTAGTTATTAATAGTAATCAATTACGGGGTCATTAGTTCATAG
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TATGCAAAGCATGCATCTCAATTAGTCAGCAACCATAGTCCCGCCCCCTAACTCCGCCCCATCCCGC

Fig. 15A

CCCTAACTCCGCCCAGTTCCGCCCATTCTCCGCCCCATGGCTGACTAATTTTTTTTTATTTATGCA
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ATCT

Fig. 15B